Amendments to the Claims:

This listing of claims will replace all prior versions and listings of the claims in the application:

Listing of Claims:

- (Currently amended) An isolated nucleic acid comprising a member 1. polynucleotide encoding a functional celllose synthase, said polynucleotide selected from the group consisting of:
 - a polynucleotide having at least 90 70% sequence identity, as (a) determined by the GAP algorithm under default parameters, to a polynucleotide selected from the group consisting of SEQ ID NOS: 25, 27 and 29, wherein the polynucleotide encodes a functional cellulose synthase;
 - a polynucleotide encoding a polypeptide selected from the group (b) consisting of SEQ ID NOS: 26, 28 and 30;
 - (c) a polynucleotide amplified from a Zea mays nucleic acid library using primers which selectively hybridize, under stringent hybridization conditions, to loci within a polynuclootide selected from the group consisting of SEQ ID NOS: 25,-27 and 29;
 - (d)a polynucleotide which selectively hybridizes, under stringent hybridization conditions and a wash in 0.1X SSC at 65°C, to a polynucleotide selected from the group consisting of SEQ ID NOS: 25, 27 and 29;
 - (e)a polynucleotide selected-from the group-consisting-of SEQ ID (c) NOS: 25, 27 and 29; and
 - (f) a polynucleotide which is complementary to a polynucleotide of (a), (d) (b), or (c), , (d), or (e); and

- (g) a polynucleotide comprising at least 25 contiguous nucleotides from a polynucleotide of (a), (b), (d), (e), or (f).
- (Currently amended) A recombinant expression cassette, comprising a member the polynucleotide of claim 1 operably linked, in sense or anti-sense orientation, to a promoter.
- (Original) A host cell comprising the recombinant expression cassette of claim 2.
- 4. (Original) A transgenic plant comprising the recombinant expression cassette of claim 2.
- 5. (Original) The transgenic plant of claim 4, wherein said plant is a monocot.
- (Original) The transgenic plant of claim 4, wherein said plant is a dicot.
- (Original) The transgenic plant of claim 4, wherein said plant is selected from the group consisting of: maize, soybean, sunflower, sorghum, canola, wheat, alfalfa, cotton, rice, barley, millet, peanut, and cocoa.
- 8. (Currently amended) A transgenic seed from the transgenic plant of claim 4.
- 9. (Currently amended) A method of modulating the level of cellulose synthase in a plant cell, comprising:

- introducing into a plant cell a recombinant expression cassette comprising a polynucleotide of claim 1 operably linked to a promoter;
 and
- (b) culturing the plant cell under plant cell growing conditions; wherein the level of cellulose synthase in said plant cell is modulated. and
- (e) expressing said-polynucleotide for a time sufficient to modulate the level of cellulose-synthase in said-plant cell.
- 10. (Currently amended) The method of claim 9, wherein the plant cell is is from a plant selected from the group consisting of: maize, soybean, sunflower, sorghum, canola, wheat, alfalfa, cotton, rice, barley, millet, peanut, and cocoa.
- 11. (Currently amended) A method of modulating the level of cellulose synthase in a plant, comprising:
 - introducing into a plant cell a recombinant expression cassette comprising a the polynucleotide of claim 1 operably linked to a promoter;
 - (b) culturing the plant cell under plant cell growing conditions; and
 - (c) regenerating a plant from said plant cell; wherein the level of cellulose synthase in said plant is modulated_and
 - (c) expressing said polynucleotide for a time sufficient to modulate the level of cellulose synthase in said plant.
- 12. (Original) The method of claim 11, wherein the plant is selected from the group consisting of: maize, soybean, sunflower, sorghum, canola, wheat, alfalfa, cotton, rice, barley, millet, peanut, and cocoa.

- 13. (Withdrawn) An isolated protein comprising a member selected from the group consisting of:
 - (a) a polypeptide of at least 20 contiguous amino acids from a polypeptide selected from the group consisting of SEQ ID NOS: 26, 28 and 30;
 - (b) a polypeptide selected from the group consisting of SEQ ID NOS: 26,28 and 30;
 - (c) a polypeptide having at least 70% sequence identity to, and having at least one epitope in common with, a polypeptide selected from the group consisting of SEQ ID NOS: 26, 28 and 30, wherein said sequence identity is determined by the GAP algorithm under default parameters; and,
 - (d) at least one polypeptide encoded by a member of claim 1.
- 14. (Withdrawn) A method of modifying expression of a cellulose synthase gene in a malze plant, comprising:
 - identifying, from a population of maize plants mutagenized with the Mu transposable element, those plants containing one or more Mu insertions within a polynucleotide of claim 1;
 - (b) selecting those plants showing modified cellulose syse gene expression.
- 15. (Withdrawn) The method of claim 14, where expression of the cellulose synthase gene is down-regulated.